

1/11

```

1  GCCACCGACA  TCCGCCGCAA  TGCTGTGTCT  CACCTCCTCT  TCCTCCTCCG  CGCCCGCTCC
61  GCTCCTTTCC  TCTCTCGCTG  ATCGACCGAG  CCCGGAATC  GCGGGCGGG  GTGGCAATGT
121  TCGCCTGAGC  GTGGTTTCTT  CGCCGCGCCG  GTCGTGGCCT  GGAAAGGTCA  AGACCAATTT
181  CTCAGTTCCCT  GCGACTGCGC  GAAAAACAA  AACCATGGTG  ACTGTTGTGG  AGGAGGTCGA
241  CCACCTTTCC  ATATATGATC  TGGACCCTAA  GTTGGAGGAA  TTCAAGGATC  ACTTCAACTA
301  TAGGATAAAA  AGATACCTCG  ACCAGAAATG  CCTGATTGAA  AAACATGAGG  GGGGCCTTGA
361  AGAATTTTCT  AAAGGCTATT  TGAAGTTTGG  GATTAATACA  GTTGATGGTG  CCACAATATA
421  TCGTGAATGG  GCGCCTGCTG  CACAAGAAGC  ACAGCTCATT  GGTGAGTTCA  ATAAGTGGAA
481  TGGTGCAAAA  CACAAGATGG  AGAAGGATAA  ATTTGGCATT  TGGTCAATCA  AGATTTTACA
541  TGTCAAATGGG  AAGCCTGCCA  TCCCTCACAA  TTCCAAGGTT  AAATTTTCGT  TTAGGCATGG
601  GGGTGGAGCA  TGGGTTGATC  GTATTCCCGC  ATGGATTTCGT  TATGCAACTT  TTGATGCCTC
661  TAAATTTGGA  GCTCCATATG  ATGGTGTACA  CTGGGATCCT  CCAGCCTGTG  AAAGGTACGT
721  GTTTAAGCAT  CCTCGACCTC  CAAAACCTGA  TGCTCCACGC  ATCTATGAGG  CTCATGTGGG
781  GATGAGTGGT  GAAGAGCCAG  AAGTAAGCAC  ATACAGAGAA  TTTGCAGACA  ATGTGTTACC
841  ACGCATACGG  GCAAATAACT  ACAACACAGT  TCAGTTAATG  GCAATCATGG  AACATTCCCTA
901  CTATGCTTCT  TTTGGGTATC  ACGTGACAAA  TTTTTTCGCA  GTCAGCAGCA  GATCAGGAAC
961  ACCAGAGGAT  CTGAAATATC  TTGTTGACAA  GGCACATAGT  TTAGGATTAC  GAGTTCTGAT
1021  GGATGTTGTC  CATAGCCATG  CGAGTAATAA  TGTGACCGAT  GGTCTAAATG  GGTATGACGT
1081  TGGACAAAAC  ACTCATGAGT  CTTATTTTCA  TACAGGAGAT  AGGGGCTACC  ATAAACTCTG
1141  GGATAGTCGT  CTGTTCAACT  ATGCCAATTG  GGAGGTCTTA  AGATTTCTTC  TTTCTAATTT
1201  GAGATATTGG  ATGGACGAAT  TCATGTTTGA  TGGCTTCCGA  TTTGATGGGG  TTACATCAAT
1261  GCTATACCAT  CACCATGGTA  TCAATAAGGG  ATTTACTGGA  AACTACAAGG  AGTATTTTCA
1321  TTTGGATACC  GATGTGGATG  CAATTGTTTA  CATGATGCTC  GCAAACCATT  TAATGCATAA
1381  ACTCTTGCCG  GAAGCAACTA  TTGTTGCTGA  AGATGTTTCG  GGCATGCCAG  TGCTTTGTCTG
1441  GCCAGTTGAT  GAAGGTGGAG  TAGGGTTTGA  CTTCCGCTG  GCAATGGCCA  TTCCTGATAG
1501  ATGGATTGAC  TACCTGAAGA  ACAAGAGGA  CCGCAAATGG  TCAATGATG  AAATAGTGCA
1561  AACTTTGACT  AACAGGAGAT  ATACAGAAAA  ATGCATTGCC  TATGCCGAGA  GCCATGATCA
1621  GTCCATTGTT  GGTGACAAGA  CTATAGCATT  TCTCTTGATG  GACAAGGAAA  TGTACACTGG
1681  CATGTCAGAC  TTGCAGCCTG  CTTACCTAC  CATCAACCGT  GGCATTGCAC  TCCAAAAGAT
1741  GATTCACTTC  ATTACGATGG  CCCTTGAGG  TGATGGCTAC  TTAAATTTTA  TGGGCAATGA
1801  GTTTGGCCAT  CCAGAATGGA  TTGACTTTCC  AAGAGAAGGC  AACAACCTGA  GCTATGATAA
1861  ATGCAGACGT  CAGTGGAGCC  TTGTCGACAC  TGATCACCTT  CGATACAAGT  ATATGAATGC
1921  ATTTGATCAA  GCAATGAATG  CACTCGAGGA  GGAATTTTCC  TTCCTGTCAT  CATCAAAGCA
1981  GATTGTTAGC  GACATGAACG  AGAAGATAA  GGTTATTGTC  TTTGAACGTG  GAGATTTGGT
2041  TTTTGTTTTT  AATTTTTCATC  CCAACAAAAC  TTACAAGGGT  TACAAAGTCG  GATGTGACTT
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2161  AGTTGGCCAT  GATGTGGATC  ACTTCACGTC  TCCCAGGGGA  ATGCCAGGAG  TACCAGAAAC
2221  AAATTTCAAC  AACCGCCCTA  ACTCATTCAA  AGTCCTTTCC  CCGCCCCGTA  CCTGTGTGGC
2281  TTACTATCGC  GTTGATGAAG  ATCGTGAAGA  GCTCAGGAGG  GGTGGAGCAG  TTGCTTCTGG
2341  AAAGATTGTT  ACAGAGTATA  TCGATGTTGA  AGCAACAAGT  GGGGAGACTA  TCTCTGGTGG
2401  CTGGAAGGGC  TCCGAGAAGG  ACGATTGTGG  CAAGAAAGGG  ATGAAGTTTG  TGTTTCGGTC
2461  TTCTGACGAA  GACTGCAAAT  GAAGCATCAG  ATTTCTTGAT  CAGGAGCAAC  TGTTGGTGCC
2521  CTTGTAATCT  GGAGATCCTG  GCTTGCTTGG  GACTTGGTTG  TGGTTCTTTA  GCAGTTGCTA
2581  TGTACCTATC  TATGATATGA  ACTTTATGTA  TAGTTCGCCT  TAAAGAAAGA  ATAAGCAGTG
2641  ATGATGTGGC  CTTAAACCTG  AGCTGCACAA  GCCTAATGTA  AAAATAAAGT  TTCAGGCTTT
2701  CATCCAGAAT  AAAACAGCTG  TTCATTTACC  ATCTCAAAA

```

Figure 1

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```

1   CTTGACTCCC CCCACTCCTC CCTCGTGCTG CTCCTCCTCG TCGCTCGGCT CGAGGCGCGG
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121 ACCGGGGGAT GCGGTCGTTC GCGGTGTCCG GCGCGAGGCT CGGGGTCTGT CGGGCGGGGG
181 GCGGCGGCGG CGGCGGGGGT GGCCCGGCGG CGCGATCCGG CGGGGTGGAC TTGCCGTCCG
241 TGCTCTTCAG GAGGAAGGAC TCCTTCTCAC GTGGCGTTGT GAGCTGCGCG GGTGCTCCTG
301 GGAAGGTGCT GGTGCCTGGC GGTGGGAGCG ACGACTTGCT GTCCTCTGCG GAACCAGACG
361 TGGAAACTCA AGAGCAACCT GAAGAATCTC AGATACCTGA TGATAATAAA GTAAACCTT
421 TTGAGGAGGA GGAAGAGATT CCAGCAGTGG CAGAAGCAAG CATAAAGGTT GTGGCTGAAG
481 ACAAACTTGA ATCTTCAGAA GTGATTCAAG ACATTGAGGA AAATGTGACT GAGGGTGTGA
541 TCAAAGATGC TGATGAACCA ACTGTGGAGG ATAAACCACG AGTTATCCCA CCACCAGGAG
601 ATGGGCAGAA GATATACCAA ATTGACCCAA TGCTGGAAGG ATTTGCGAAC CATCTTGACT
661 ACCGATACAG TGAATACAAG AGAATGCGTG CAGCTATTGA CCAACATGAA GGTGGCTTGG
721 ATGCATTTTC TCGTGGTTAC GAAAAGCTTG GATTCACCCG CAGCGCTGAA GGCATTACCT
781 ACCGAGAATG GGCACCTGGA GCACAGTCTG CAGCATTAGT AGGTGACTTC AACAATTGGA
841 ACCCAAATGC AGATACTATG ACCAGAAATG AGTATGGTGT TTGGGAGATT TCCCTGCCTA
901 ACAATGCTGA TGGATCCCCCT GCTATTCCTC ATGGCTCACG TGTAAGATT CGGATGGATA
961 CACCATCTGG CGTAAAGGAT TCAATTCTCT CCTGGATTAA GTTTGCTGTG CAGGCTCCAG
1021 GTGAAATACC GTACAACGGT ATATATTATG ATCCACCTGA AGAAGAAAAA TATGTATTCC
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1141 GTAGCCCGGA ACCGAAGATA AACACATATG CTAATTTTAG GGATGAGGTG CTACCAAGAA
1201 TTAAAAAGCT TGGGTACAAT GCTGTACAGA TAATGGCAAT CCAGGAGCAC TCTTATTACG
1261 CAAGCTTTGG GTATCATGTT ACTAACTTCT TTGCGCCAAG TAGCCGTTTC GGAACCCAG
1321 AAGACTTGAA ATCTCTGATT GATAAAGCTC ACGAGCTTGG TTTGCTTGTA CTTATGGATA
1381 TTGTTACAG TCATGCATCA AACAATACCC TGGATGGTTT GAATGGTTT GATGGTACTG
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1501 TCAACTATGG GAGTTGGGAA GTTTTAAGAT ATTTACTGTC GAATGCAAGG TGGTGGCTTG
1561 AAGAATACAA GTTTGATGGG TTTGATTTG ATGGGGTGAC CTCCATGATG TATACTCATC
1621 ATGGTTTACA GGTGGCATT TACTGGCACT ATGGCGAATA TTTTGGATTT GCTACTGATG
1681 TTGATGCAGT AGTTTACTTG ATGCTGGTGA ACGATCTAAT TCATGGGCTT TATCCTGAGG
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1921 GGTGGTCAGA GAAGTGTGTT ACTTATGCAG AAAGTCATGA CCAAGCACTA GTTGGTGACA
1981 AGACTATTGC ATTCTGGTTG ATGGATAAGG ATATGTATGA TTTTATGGCT CTAGACAGAC
2041 CTTCAACACC TCGCATTGAT CGTGGGATAG CATTACATAA AATGATTAGG CTTGTCACCA
2101 TGGGCTTAGG AGGCGAAGGC TATCTTAATT TCATGGGAAA TGAGTTTGGG CATCCTGAAT
2161 GGATAGATTT CCAAGAGGC CCGCAAAGTC TTCCAAATGG CTCGGTCCTC CCAGGAAACA
2221 ACTACAGTTT TGATAAATGC CGTCGTAGAT TTGACCTTGG AGATGCAGAT TATCTTAGAT
2281 ATCATGGTAT GCAAGAGTTT GATCAGGCCA TGCAGCATCT TGAGGAAAAA TATGGATTCA
2341 TGACATCTGA GCACCAGTAT ATATCGCGCA AACACGAGGA GGATAAGGTG ATCATCTTCG
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2641 TTACAGAGGA CTAATGATCA GCTCTGATCA TTGGGGGAAC AACTCAAGGG AGTTGGTGGT
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2881 AGCTATACCA AACCCATCCT ATGTTGCGCA TTCGCTGTAG TTTTGTACAT AACGATATCG
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Figure 2

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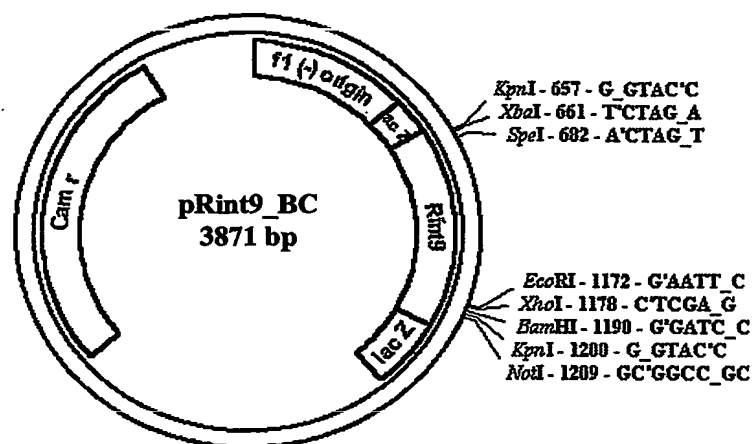
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121  GGGCGAGATG  GCGGCGCCGG  CGTCTGCGGT  TCCC GGAGC  GCGGCGGGGC  TACGGGCGGG
181  GGCCGTGCGG  TTCCCCGTGC  CAGCCGGGGC  CCGGAGCTGG  CGTGCGGGCG  CGGAGCTCCC
241  GACGTGCGGG  TCGCTGCTCT  CCGGCCGGAG  ATTCCCCGGT  GCCGTTGCGG  TGGGGGGTTC
301  CGGGGGGCGC  GTGGCCGTGC  GCGCGGCGGG  CGCGTCAGGG  GAGGTGATGA  TCCCCGAGGG
361  CGAGAGCGAC  GGGATGCCGG  TTTTCAGCAG  TTCAGACGAT  CTGCAGTTGC  CAGCCTTAGA
421  TGATGAATTA  AGCACGGAGG  TTGGAGCTGA  AGTTGAGATT  GAGTCATCTG  GAGCAAGTGA
481  CGTTGAAGGC  GTGAAGAGAG  TGGTTGAAGA  ATTAGCTGCT  GAGCAGAAAC  CACGAGTTGT
541  CCCACCAACA  GGAGATGGGC  AAAAAATATT  CCAGATGGAC  TCTATGCTTA  ATGGCTATAA
601  GTACCATCTT  GAATATCGAT  ATAGCCTATA  TAGGAGACTG  CGTTCAGACA  TTGATCAGTA
661  TGAAGGAGGA  CTGGAACAT  TTTCTCGCGG  TTATGAGAAG  TTTGGATTTA  ATCACAGTGC
721  TGAAGGTGTC  ACTTATCGAG  AATGGGCTCC  CGGGGCACAT  TCTGCAGCAT  TAGTAGGTGA
781  CTTCAACAAT  TGGAATCCAA  ATGCAGACCG  CATGAGCAAA  AATGAGTTTG  GTGTTTGGGA
841  GATTTTTCTG  CCTAACAAAT  CTGATGGCTC  ATCTCCTATT  CCACATGGCT  CACGTGTAAG
901  GGTGCGAATG  GAAACTCCAT  CTGGTATAAA  GGATTCTATT  CCTGCCTGGA  TCAAGTACTC
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1141  GGTGCTTCCA  AGAATCAAAA  AGCTTGGATA  CAATGCAGTG  CAAATAATGG  CAATTCAAGA
1201  GCATGCATAT  TATGGAAGCT  TTGGGTACCA  TGTCACCAAT  TTCTTTGCAC  CAAGTAGTCG
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1381  TTTTGATGGT  ACAGATACGC  ATTACTTTCA  TAGTGGTTCA  CGCGGCCATC  ATTGGATGTG
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1981  GGCTCTGGAC  AGACCGGCAA  CACCTAGCAT  TGATCGTGGA  ATAGCATTGC  ATAAAATGAT
2041  TAGACTTATC  ACAATGGGGT  TAGGAGGAGA  AGGCTATCTT  AACTTTATGG  GAAATGAGTT
2101  CGGACATCCT  GAATGGATTG  ATTTTCCAAG  AGCTCCACAA  GTACTTCCAA  ATGGTAAATT
2161  CATCCAGGG  AATAACAACA  GTTATGATAA  ATGCCGTCGA  AGATTTGACC  TGGGTGATGC
2221  GGAATATCTT  AGGTATCGTG  GCATGCTAGA  GTTTGACCGC  GCGATGCAGT  CTCTCGAGGA
2281  AAAATATGGG  TTCATGACAT  CAGACCACCA  GTACATATCT  CGAAAGCATG  AAGAGGATAA
2341  GATGATTATA  TTTGAGAAGG  GAGATCTGGT  ATTTGTGTTT  AACTTCCATT  GGAGTAACAG
2401  CTATTTTGAC  TACCGTGTG  GTTGTTTAAA  GCCAGGGAAA  TATAAGGTGG  TCTTGACTC
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2641  GCTAGTGCGA  AGGAGCAAGA  AAAACTAGTT  GCCAGCAATC  TGTGAACGGC  TTTCTTAGGT
2701  TCTGCTTCGA  TGAATGCCGG  ATAGACTAGA  CAGCTTGCTT  TTGTGCTTTG  CGCTCCCAAT
2761  TTGTAGTTTT  AGTTTGTGAG  GGAAAGAAAC  GTTTATTTGT  AATTATCTAT  GGCTGTCGAA
2821  CGGCGACGAA  ACCATGAACC  CCGTATATTT  GTTGGTACCG  TTCGAAGTGC  CAGTTATACA
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Figure 3

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**Figure 4**

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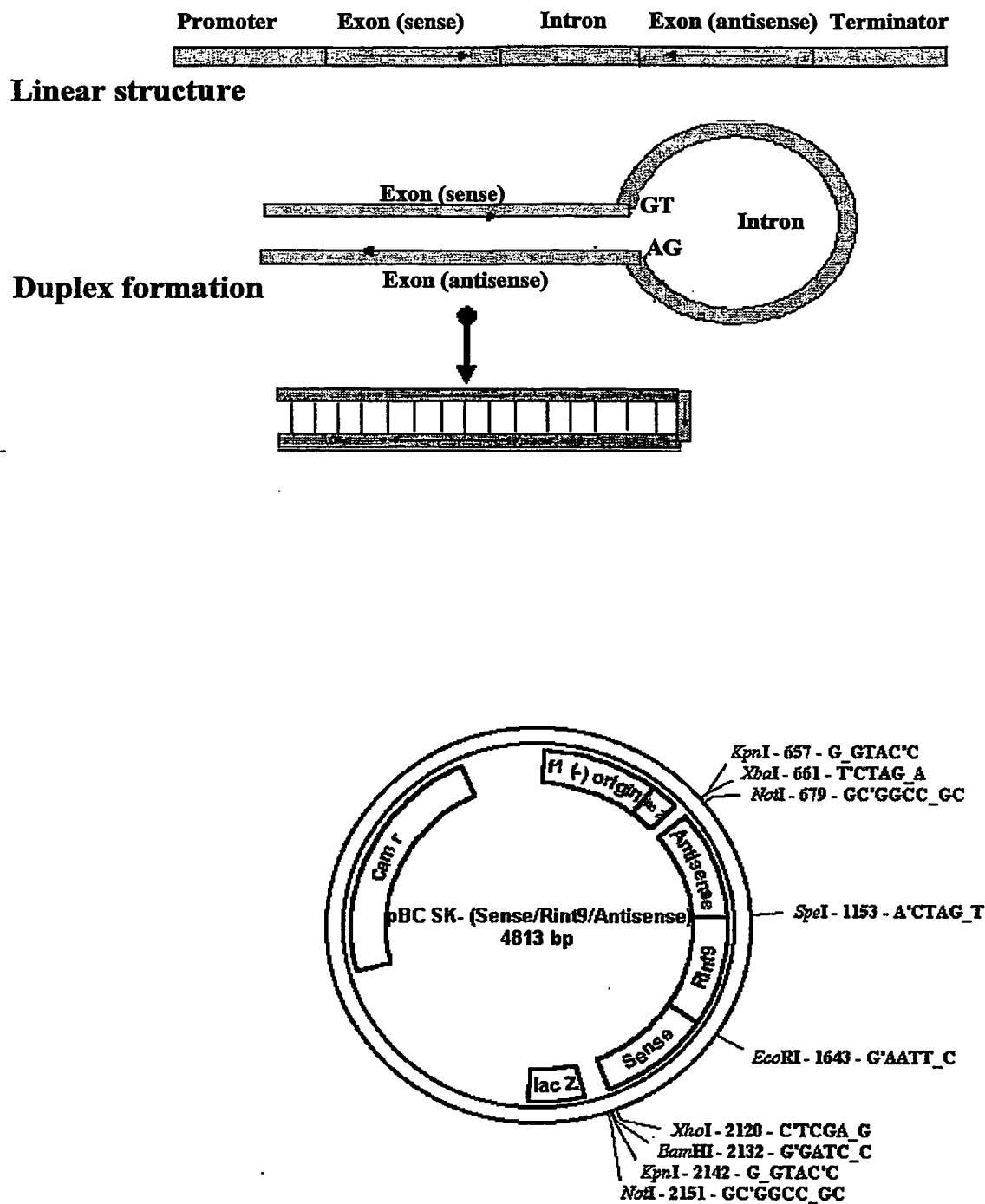
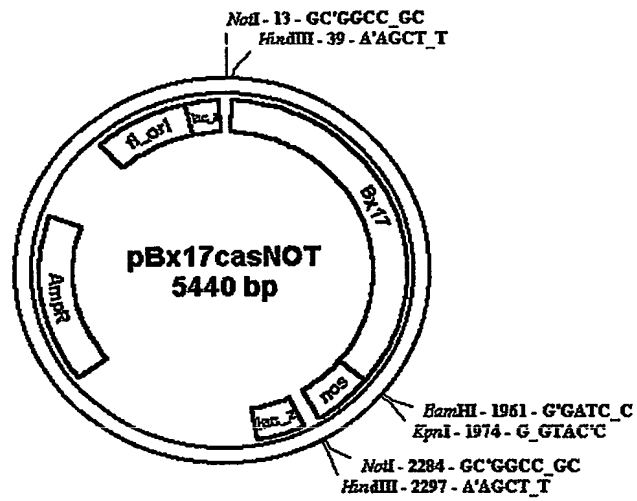


Figure 5

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**Figure 6**

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```

82  GCGCGGGGGTTGCCGGGGGATCCGATCCGGCTGCG.GTGCGGGCGAGATG 130
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55  gcgcggcatttgcgggcgga.gggatctgcgcgcgagtgcggtcgggcgag 103

131  GCGGC.....GCCGGCGTCTGCGGTTCCCGGGA 158
   ||||| ||||| ||||| ||||| ||||| |||||
104  gcggcggggggagcacgcaccgggggatggcgctcggtcgcggtgtcc.ggc 152

159  GCGCGGGCGGGGCTACGGGCGGGGGCCGTGCGGTTCCCCGTGCCAGCCGGG 208
   ||||| ||||| ||||| ||||| ||||| ||||| |||||
153  gcgaggctcggggtcggtgcgggcggggggcg...cggcggcgggcgggg 198

209  GCGCGGAGCTGGCGTGCGGCGGCGGAGCTCCCGACGTGCGGTCGCTGCT 258
   ||||| ||||| ||||| ||||| ||||| ||||| |||||
199  gtggcccgggcggcgcgatccggcgggg....tggacttgccgtcggtgct 244

259  CTCCGGCCGGAGATTCCCCGGTGCCGTTTCGCGTGGGGGGTTCCGGGGGGC 308
   ||||| ||||| ||||| ||||| ||||| ||||| |||||
245  cttcaggagga.....aggactccttctcacgtggcggt..... 278

309  GCGTGCCGTGCGCGCGGGCGGCGCTCAGGGGAGGTGATGATCCCCGAG 358
   ||||| ||||| ||||| ||||| ||||| ||||| |||||
279  .....gtgagctgcgcgggtgctcctgggaaggtgctggtgcctggc 320

359  GCGGAGAGCGACGGGATGCCGGTTTCAGCAGGTTTCAGACG..... 398
   ||||| ||||| ||||| ||||| ||||| ||||| |||||
321  ggtgggagcgacgacttgctgtcctctgcggaaccagacgtggaaactca 370

399  .....ATCTGCAGTTGCC.....AGCCT 416
   ||||| ||||| ||||| ||||| ||||| |||||
371  agagcaacctgaagaatct.cagataacctgatgataataaaagtaaacct 419

417  T.....AGATGATGAATTAAGCACGGAGGT 441
   ||||| ||||| ||||| ||||| ||||| |||||
420  tttgaggaggaggaagagattccagcagtggcagaagcaagcataaagggt 469

442  TGGAGCTGAAGTTGAGATTGAGTCATC.....TGGAG 473
   ||||| ||||| ||||| ||||| ||||| |||||
470  tgtggctgaagacaaacttgaatcttcagaagtgattcaagacattgagg 519

474  CAAGTGACGTTGAAGGCGTGAAGAGAGTGGTTGAAGAATTAGCTGCTGAG 523
   ||||| ||||| ||||| ||||| ||||| |||||
520  aaaatgtgactgaggggtgtgatcaaagatgctgatgaaccaactgtggag 569

524  CAGAAACCACGAGTTGTCCCACCAACAGGAGATGGGCAAAAAATATTCCA 573
   ||||| ||||| ||||| ||||| ||||| ||||| |||||
570  gataaaccacgagttatcccaccaccaggagatgggcagaagatatacca 619

574  GATGGACTCTATGCTTAATGGCTATAAGTACCATCTTGAATATCGATATA 623
   ||||| ||||| ||||| ||||| ||||| |||||
620  aattgacccaatgctggaaggatttcggaaccatcttgactaccgataca 669

624  GCCTATATAGGAGACTGCGTTTCAGACATTGATCAGTATGAAGGAGGACTG 673
   ||||| ||||| ||||| ||||| ||||| |||||
670  gtgaatacaagagaatgcgtgcagctattgaccaacatgaagggtggcttg 719

```

Figure 7

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674 GAAACATTTTCTCGCGGTTATGAGAAGTTTGGATTTAATCACAGTGCTGA 723
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720 gatgcattttctcgtggttacgaaaagcttggattcaccgcgagcgctga 769
    .
724 AGGTGTCACCTATCGAGAATGGGCTCCCGGGGCACATTCGTCAGCATTAG 773
    ||| | ||| ||||| ||||| ||||| ||||| ||||| |||||
770 aggcattacctaccgagaatgggcacctggagcacagtctgcagcattag 819
    .
774 TAGGTGACTTCAACAATTGGAATCCAAATGCAGACCGCATGAGCAAAAAT 823
    ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
820 taggtgacttcaacaattggaacccaaatgcagatactatgaccagaaat 869
    .
824 GAGTTTGGTGTGTTGGGAGATTTTCTGCCTAACAATGCTGATGGCTCATC 873
    ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
870 gagtatggtggttgggagatttccctgcctaacaatgctgatggatcccc 919
    .
874 TCCTATTCCACATGGCTCACGTGTAAAGGTGCGAATGGAACTCCATCTG 923
    | ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
920 tgctattcctcatggctcacgtgtaaagattcggatggatacaccatctg 969
    .
924 GTATAAAGGATTCTATTCTGCCTGGATCAAGTACTCTGTGCAGGCCGCA 973
    | ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
970 gcgtaaaggattcaattcctgcctggattaagtttgctgtgcagggtcca 1019
    .
974 GGAGAAATCCCATACAATGGAATATATTATGATCCTCCTGAAGAGGAGAA 1023
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1124 GCAAACTTTAGGGATGAGGTGCTTCCAAGAATCAAAAAGCTTGGATACAA 1173
    || || ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
1170 gctaatttttagggatgaggtgctaccaagaattaaaaagcttgggtacaa 1219
    .
1174 TGCAGTGCAAATAATGGCAATTCAAGAGCATGCATATTATGGAAGCTTTG 1223
    ||| || ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
1220 tgctgtacagataatggcaatccaggagcactcttattacgcaagctttg 1269
    .
1224 GGTACCATGTCACCAATTCCTTTGCACCAAGTAGTCGTTTCGGGACCCCA 1273
    ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
1270 ggtatcatgttactaacttctttgcgccaagtagccgtttcggaacccca 1319
    .
1274 GAAGATTTAAAGTCATTGATTGATAAAGCTCATGAGCTTGGTTTAGTTGT 1323
    ||||| ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
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    .
1324 GCTCATGGATGTTGTTTCACAGCCATGCGTCAAATAATACCTTAGATGGGT 1373
    || ||||| ||||| ||||| ||||| ||||| ||||| ||||| |||||
1370 acttatggatattgttcacagtcatgcatcaaacaataccctggatgggt 1419

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Figure 7



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|||  
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1474 AGTTCTAAGATTTCTACTATCCAATGCAAGATGGTGGCTCGAGGAGTATA 1523  
|||||  
1520 agttttaagatatttactgtcgaatgcaagggtgtggcttgaagaataca 1569  
1524 AGTTTGATGGTTTCAGATTTGACGGTGTAACTCAATGATGTACACTCAT 1573  
|||||  
1570 agtttgatgggtttcgatttgatgggggtgacctccatgatgtatactcat 1619  
1574 CATGGATTACAAGTAGCATTTTACGGGGAAC TACAGTGAATAC TTTGGATT 1623  
|||||  
1620 catggtttacagggtggcatttactggcaactatggcgaatattttggatt 1669  
1624 TGCCACTGATGCTGATGCAGTAGTTTACTTGATGCTGGTAAATGATT TAA 1673  
|||  
1670 tgctactgatgttgatgcagtagtttacttgatgctggtgaacgatctaa 1719  
1674 TTCATGGACTTTATCCTGAGGCCATAACCATCGGTGAAGATGTCAGTGGA 1723  
|||||  
1720 ttcatgggctttatcctgaggctgtagccattgggtgaagatgtcagcggg 1769  
1724 ATGCCTACATTTGCCCTTCCTGTTCAAGATGGTGGGGTTGGTTTTGATTA 1773  
|||||  
1770 atgcccacattttgtattcctgttcaagatgggtggtggttttgacta 1819  
1774 TCGCCTTCATATGGCTGTTCTGACAAATGGATTGAACTCCTCAAGCAAA 1823  
|||  
1820 tcgtttgcatatggctgtaccggacaaatggatcgaactcctcaagcaaa 1869  
1824 GTGATGAATCTTGGAAGATGGGTGATATTGTGCACACACTGACTAACAGA 1873  
|||||  
1870 gtgacgaatattggaaaatgggtgatatcggtgcacaccctaacgaataga 1919  
1874 AGGTGGTCAGAGAAGTGTGTTACTTATGCTGAAAGTCATGATCAAGCACT 1923  
|||||  
1920 aggtggtcagagaagtggttacttatgcagaaagtcacgaccaagcact 1969  
1924 AGTTGGTGACAAACTATTGCATTCTGGTTGATGGACAAGGATATGTATG 1973  
|||||  
1970 agttggtgacaagactattgcattctggttgatggataaggatatgtatg 2019  
1974 ATTTTATGGCTCTGGACAGACCGGCAACACCTAGCATTGATCGTGGAATA 2023  
|||||  
2020 attttatggctctagacagaccttcaacacctcgattgatcggtgggata 2069  
2024 GCATTGCATAAAATGATTAGACTTATCACAATGGGGTTAGGAGGAGAAGG 2073  
|||||  
2070 gcattacataaaatgattaggcttgccacctgagggttaggagggcgaagg 2119

Figure 7

### Figure 7

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## riceSBEIIaIR.seq

```

1      CTCGAGTCTA  GATCGCGTC  G GTTGTTTA  AA GCCTGGA  AAG TACAAG  ATTGT
56     GTTGGA CTC  AGACGATGGC  CTCTTTGGT  G GATTCAGT  CG GCTTGAT  CATGA
111    TGCTGAGT A  CTTCACTGC  TGACTGGCCG  CATGACAAC  A GACCATGT  TCATT
166    CTCGGTG TA  CACCCCAA  G CAGAACCGC  CGTCGTGTAT  GCACTTACA  GAGGA
221    CTAATG ATC  AGCTCTG  AT CATTTGGG  G AACAACTCA  AGGGAGTTGG  TGGTA
276    ATGAC GCCG  GAATAC  AAC TCAAGTG  AA AGGTGAAA  A GAAAGGCTGC  CCTGA
331    CGAT GTGAT  TTGAG  GGGC TTGTGT  TTC ATCGCCA  AT GCCAGGAAGA  TGAGG
386    TAG AAAAGC  CTAC  TGATG  AGCTC  CTGT TTTCGA  GTG ACTCGTGAAG  GAAAT
441    AG ACCAGGG  TGA  ACGGCT  TTTT  TCAGA  GCTAT  ACCA AACCCATCCT  ATGTT
496    G CGCATTCG  CT  GTAGTTT  TGT  ACATAA  CGAT  ATCGG  TTGGCATTG  TATGT
551    TTATGAATA  A  TCTGTTCG  AC  AGAAATG  TTT  TTCTCC  TTGTAAGTAG  TGAA
606    TTC

```

## riceSBEIIbIR.seq

```

1      CTCGAGTCTA  GNNNNNNNN  N NNNNNNNN  NN NNNNNNN  NNN NNNNNNN  NNNNN
56     NNNNNNNNNN  NNNNNNNNNN  NNNNNNNNNN  N NNNNNNNN  NN NNNNNNN  NNNNN
111    NNNNNNNNN  N  NNNNNNNNN  NNNNNNNNNN  NNNNNNNNN  N NNNNNNNN  NNNNN
166    NNNNNNNN  NN  NNNNNNNN  N NNNNNNNNN  NNNNNNNNNG  CTCCAGCGG  AATGA
221    GAACAC  CAA  GAGGCAG  CA  TGCAAGTG  T  GTGCGGCTG  CTAGTGCGAA  GGAGC
276    AAGAA  AAAC  TAGTTG  CCA  GCAATCT  GT  GAACGGCT  T  TCCTAGGTTC  TGCTT
331    CGAT  GAATG  CCGGA  TAGA  CTAGAC  ANN  NNNNNNN  NN  NNNNNNNNNN  NNNNN
386    NNT  TGTAGT  TTTA  GTTTG  TGAGG  GAAA  GAAACG  TTT  ATTTGTAATT  ATCTG
441    TG  GCTGTCG  AAC  GGCGAC  GAAA  CCATG  AACCC  CGTA  TATTTGTTGG  TACCG
496    T  TCGAACTG  CC  AGTTATA  CAT  AGTTCT  GCAC  TTCTG  TACATCTTGT  GATGC
551    TACTAGTGA  A  TTC

```

## riceSBEIIR.seq

```

1      CTCGAGTCTN  NNNNNNNNNN  NNNNNNNNNN  NNNNNNNNNN  NNNNNNNNNN  NNNNN
56     NNNNNNNNNN  NNNNNNNNNN  NNNNNNNNNN  NNNNNNNNNN  NNNNNNNNNN  NNNNN
111    NNNNNNNNNN  NNNNNNNNNN  NNNNNNNNNN  NNNNNNNNNN  NNNNNNNNNN  NNNNN
166    NNNNNNNNNN  NNNNNNNNNN  NNNNNNNNNN  NNNNNNNNNN  NNNNNNNNNN  NNNNN
221    NNNNNNNNNN  NNNNNNNNNN  NNNNNNNNNN  NNNNNNNNNN  NNNNNNNNNN  NNNNN
276    NNNNNNNNNN  NNNNNNNNNN  NNNNNNNNNN  NNNNNNNNNN  NNNNNNNNNN  NNNNN
331    NNNNNNNNNN  NNNNNNNNNN  NNNNNNNNNN  NNNNNNNNNN  NNNNNNNNNN  NNNNN
386    NNNNNNNNNN  NNNNNNNNNN  NNNNNNNNNN  NNNNNNNNNN  NNNNNNNNNN  NNNNN
441    NNNNNNNNNN  NNNNNNNNNN  NNNNNNNNNN  NNNNNNNNNN  NNNNNNNNNN  NNNNN
496    NNNNNNNNNN  NNNNNNNAGC  ATCAGATTTC  TTGATCAGGA  GCAACTGTTG  GTGCC
551    CTTGTAAACT  AGTGAATTC

```

Figure 8